

SEQUENCE LISTING

<110> Basson, Craig

<120> Transcription Factors that Regulate Normal and Malignant Cell Growth

<130> 955-12P

<160> 15

<170> PatentIn version 3.0

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<211> 349

<212> PRT

<213> Homo sapiens

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Gly Ala Pro Ser Lys Ser Pro Ser Ser Pro Gln Ala Ala Phe Thr Gln
 35 40 45

Gln Gly Met Glu Gly Ile Lys Val Phe Leu His Glu Arg Glu Leu Trp
 50 55 60

Leu Lys Phe His Glu Val Gly Thr Glu Met Ile Ile Thr Lys Ala Gly
 65 70 75 80

Arg Arg Met Phe Pro Ser Tyr Lys Val Lys Val Thr Gly Leu Asn Pro
 85 90 95

Lys Thr Lys Tyr Ile Leu Leu Met Asp Ile Val Pro Ala Asp Asp His
 100 105 110

Arg Tyr Lys Phe Ala Asp Asn Lys Trp Ser Val Thr Gly Lys Ala Glu
 115 120 125

Pro Ala Met Pro Gly Arg Leu Tyr Val His Pro Asp Ser Pro Ala Thr
 130 135 140

Gly Ala His Trp Met Arg Gln Leu Val Ser Phe Gln Lys Leu Lys Leu
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Thr Asn Asn His Leu Asp Pro Phe Gly His Ile Ile Leu Asn Ser Met
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His Lys Tyr Gln Pro Arg Leu His Ile Val Lys Ala Asp Glu Asn Asn
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Gly Phe Gly Ser Lys Asn Thr Ala Phe Cys Thr His Val Phe Pro Glu
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Thr Ala Phe Ile Ala Val Thr Ser Tyr Gln Asn His Lys Ile Thr Gln
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 Leu Lys Ile Glu Asn Asn Pro Phe Ala Lys Gly Phe Arg Gly Ser Asp
 225 230 235 240
 Asp Met Glu Leu His Arg Met Ser Arg Met Gln Ser Lys Glu Tyr Pro
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 260 265 270
 Pro Phe Ser Ser Glu Ser Arg Ala Leu Ser Thr Ser Ser Asn Leu Gly
 275 280 285
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 290 295 300
 Leu Pro Pro Pro Asn Pro Tyr Pro Leu Pro Gln Glu His Ser Gln Ile
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 Gly Ala Pro Ser Lys Ser Pro Ser Ser Pro Gln Ala Ala Phe Thr Gln
 35 40 45
 Gln Gly Met Glu Gly Ile Lys Val Phe Leu His Glu Arg Glu Leu Trp
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 Leu Lys Phe His Glu Val Gly Thr Glu Met Ile Ile Thr Lys Ala Gly
 65 70 75 80
 Arg Arg Met Phe Pro Ser Tyr Lys Val Lys Val Thr Gly Leu Asn Pro
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 Lys Thr Lys Tyr Ile Leu Leu Met Asp Ile Val Pro Ala Asp Asp His
 100 105 110
 Arg Tyr Lys Phe Ala Asp Asn Lys Trp Ser Val Thr Gly Lys Ala Glu
 115 120 125
 Pro Ala Met Pro Gly Arg Leu Tyr Val His Pro Asp Ser Pro Ala Thr
 130 135 140
 Gly Ala His Trp Met Arg Gln Leu Val Ser Phe Gln Lys Leu Lys Leu
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 Thr Asn Asn His Leu Asp Pro Phe Gly His Ile Ile Leu Asn Ser Met
 165 170 175
 His Lys Tyr Gln Pro Arg Leu His Ile Val Lys Ala Asp Glu Asn Asn
 180 185 190
 Gly Phe Gly Ser Lys Asn Thr Ala Phe Cys Thr His Val Phe Pro Glu
 195 200 205

Thr Ala Phe Ile Ala Val Thr Ser Tyr Gln Asn His Lys Ile Thr Gln
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 Leu Lys Ile Glu Asn Asn Pro Phe Ala Lys Gly Phe Arg Gly Ser Asp
 225 230 235 240
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 Val Val Pro Arg Ser Thr Val Arg Gln Lys Val Ala Ser Asn His Ser
 260 265 270
 Pro Phe Ser Ser Glu Ser Arg Ala Leu Ser Thr Ser Ser Asn Leu Gly
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 Pro Met Asp Arg Leu Pro Tyr Gln His Phe Ser Ala His Phe Thr Ser
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 Gly Pro Leu Val Pro Arg Leu Ala Gly Met Ala Asn His Gly Ser Pro
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 Gln Leu Gly Glu Gly Met Phe Gln His Gln Thr Ser Val Ala His Gln
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 Pro Val Val Arg Gln Cys Gly Pro Gln Thr Gly Leu Gln Ser Pro Gly
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 Thr Leu Gln Pro Pro Glu Phe Leu Tyr Ser His Gly Val Pro Arg Thr
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<210> 8
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